

INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)

Applicant's or agent's file reference 12367500/TDO/FT	FOR FURTHER ACTION	See Notification of Transmittal of International Preliminary Examination Report (Form PCT/IPEA/416).			
International Application No.	International Filing Date (day/month/year)	е	Priority Date (day/month/year)		
PCT/AU2003/001497	13 November 2003		13 November 2002		
International Patent Classification (IPC) or national classification and IPC					
Int. Cl. 7 C12N 5/00, 5/06, 5/08					
Applicant					
MONOQUANT PTY LTD et al					
1. This international preliminary examination report has been prepared by this International Preliminary Examining Authority and					
is transmitted to the applicant according	g to Article 36.				
2. This REPORT consists of a total of 5	sheets, including this co	over sheet.			
This report is also accompanied by ANNEXES, i.e., sheets of the description, claims and/or drawings which have been amended and are the basis for this report and/or sheets containing rectifications made before this Authority (see Rule 70.16 and Section 607 of the Administrative Instructions under the PCT).					
These annexes consist of a total o	of sheet(s).				
3. This report contains indications relating	to the following items:				
I X Basis of the report					
II Priority	II Priority				
III Non-establishment of opi	inion with regard to nove	lty, inventive step ar	nd industrial applicability		
IV Lack of unity of invention	Lack of unity of invention				
	ment under Article 35(2) with regard to novelty, inventive step or industrial applicability; explanations supporting such statement				
VI Certain documents cited	Certain documents cited				
VII Certain defects in the inte	rnational application				
VIII X Certain observations on the	VIII X Certain observations on the international application				
Date of submission of the demand Date of completion of the report					
31 May 2004		21 February 2005			
Name and mailing address of the IPEA/AU	· A	Authorized Officer			
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1.	Basis of the repo	rt				
1.	With regard to the ele	th regard to the elements of the international application:*				
	X the internationa	the international application as originally filed.				
	the description,	pages , as originally filed,				
		pages, filed with the demand,				
		pages, received on with the letter of				
	the claims,	pages, as originally filed,				
		pages, as amended (together with any statement) under Article 19,				
		pages , filed with the demand,				
		pages, received on with the letter of				
	the drawings,	pages, as originally filed,				
		pages, filed with the demand,				
		pages, received on with the letter of				
	the sequence lis	the sequence listing part of the description:				
		pages , as originally filed				
		pages, filed with the demand				
		pages, received on with the letter of				
2.	which the international	With regard to the language, all the elements marked above were available or furnished to this Authority in the language in which the international application was filed, unless otherwise indicated under this item. These elements were available or furnished to this Authority in the following language which is:				
	the language of	a translation furnished for the purposes of international search (under Rule 23.1(b)).				
	the language of	publication of the international application (under Rule 48.3(b)).				
	the language of and/or 55.3).	the translation furnished for the purposes of international preliminary examination (under Rules 55.2				
3.		ith regard to any nucleotide and/or amino acid sequence disclosed in the international application, the international preliminary examination was carried out on the basis of the sequence listing:				
	contained in the	international application in written form.				
	filed together wi	ith the international application in computer readable form.				
	furnished subsec	quently to this Authority in written form.				
	furnished subsec	furnished subsequently to this Authority in computer readable form.				
		The statement that the subsequently furnished written sequence listing does not go beyond the disclosure in the international application as filed has been furnished.				
	The statement the been furnished	at the information recorded in computer readable form is identical to the written sequence listing has				
4.	The amendment	s have resulted in the cancellation of:				
	the desc	cription, pages				
	the clair	ms, Nos.				
	the drav	wings, sheets/fig.				
<u>5</u> .		been established as if (some of) the amendments had not been made, since they have been considered to isclosure as filed, as indicated in the Supplemental Box (Rule 70.2(c)).**				
*		Replacement sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to this report since they do not contain amendments (Rules 70.16 and 70.17).				
**	Any replacement sheet	containing such amendments must be referred to under item 1 and annexed to this report				



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V. Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement

1. Statement				
	Novelty (N)	Claims 13, 14 and 26-29	YES	
		Claims 1-12, 15-25, 30 and 31	NO	
	Inventive step (IS)	Claims 13, 14, 28 and 29	YES	
		Claims 1-12, 15-27, 30 and 31	NO	
	Industrial applicability (IA)	Claims 1-31	YES	
		Claims	NO	

2. Citations and explanations (Rule 70.7)

The invention relates to methods of detecting clonal populations of cells by co-localising a subject nucleic acid region from a population of cells in a biological sample and assessing the level of the co-localised nucleic acid relative to a background level, where a higher comparative level is indicative of a clonal population of cells. In particular the description exemplifies assessing the mitochondrial D-loop region using denaturing gradient gel electrophoresis (DGGE).

The following documents identified in the International Search Report have been considered for the purposes of this report:

- D1 Cancer Research 61, 7015-19
- D2 Biochem Biophys Res Comm 199, 511-18
- D3 Diagnost Mol Path 6, 140-6
- D4 Diagnost Mol Path 6, 71-77
- D5 J Surg Res 85, 311-16
- D6 Leukemia 8, 946-52
- D7 J Neuropath Exp Neurol 61, 396-402
- D8 J Clin Path: Mol Path 53, 150-54
- D9 Int J Parasitol 32, 27-38
- D10 WO 2002 088388
- D12 US 2002/0004201

Novelty and Inventive Step

Documents D1 and D10 are not directly relevant to the novelty or inventive step of the claims. Although D1 discloses mutations in the D loop in cancers and use of this region to identify the origin and clonality of cancer cells, it does not disclose co-localising nucleic acid regions. It simply discloses amplification and sequencing. D10 discloses a method of characterising mutation loads and heterozygosity, but does not suggest that such a method can also be used to characterise the extent of clonality.

D2 discloses a method of detecting and assessing clonality using PCR to amplify target regions and DGGE to identify, quantify and compare levels of different variants within a population of cells. The target regions are an intron in the HPRT gene and a region in the PGK gene and the clonality is associated with cancer. As such D2 deprives claims 1-3, 11, 15-18, 26, 30 and 31 of novelty and an inventive step.

Continued in supplemental box.



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VIII. Certain observations on the international application

The following observations on the clarity of the claims, description, and drawings or on the question whether the claims are fully supported by the description, are made:

Claims 1-14 and 17-29 are not fully supported by the description. The description discloses methods of co-localisation that separate and co-localise variants on the basis of differing molecular sizes of the variants arising from nucleic acid sequence variations within a distinctive nucleic acid region. The relative amounts of the separated variants are then used to generate a profile that is compared with a background profile of the region.

In contrast the claims do not specify that the method requires separation on the basis of sequence variation and molecular size followed by preparation of a profile that reflects relative amounts of sequence variants. As such the claims do not recite all of the essential features that define the applicant's invention.



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Supplemental Box

(To be used when the space in any of the preceding boxes is not sufficient)

Continuation of Box V2

D3-D6 disclose the use of PCR and denaturing PAGE to characterise clonality in B and T cell cancers, in particular lymphomas and leukaemias. As such D3-D6 deprive claims 1-5, 8, 9, 11, 15-20, 23, 24, 26, 30 and 31 of novelty and an inventive step.

D8 also discloses the use of PCR and PAGE to characterise clonality in T cell populations. However D8 is not restricted to T cell cancers and discusses other T cell conditions. As such, D8 is not only relevant to the novelty and inventive step of claims 1-5, 8, 9, 11, 15-20, 23, 24, 26, 30 and 31, it also deprives claims 6 and 21 of novelty and an inventive step.

D7 discloses the use of PCR and electrophoresis to identify clonality in gliomas. In particular the citation assesses differences in microsatellite sequences. As such D7 deprives claims 1-3, 11, 12, 15-18, 26, 27, 30 and 31 of novelty and an inventive step.

In addition, D7 also deprives claims 4, 5 and 19 of an inventive step. Although these claims relate to methods of essing clonality in B and T cell cancers and D7 relates to gliomas, the discussion in D7 discloses the use of such a such property of the control o

D9 discloses analysis of clonality in populations of the single-celled parasiste *Toxoplasma gondii* using PCR and electrophoresis. The regions that is analysed is a microsatellite region. As such D9 deprives claims 1, 2, 6, 10-12, 15-17, 21, 25-27, 30 and 31 of novelty and an inventive step.

D11 discloses analysis of loss of heterozygositiy and assessment of clonality in a range of cells and populations, including populations comprising cancerous cells and populations of microorganisms. The analysis includes PCR amplification and binding of probes to specific variants. The selected populations of variants may then be separated or co-localised as a consequence of affinity binding of the probes to binding matrices or as a consequence of separation of probes based on differing molecular weight or size. As such D11 deprives claims 1-11, 15-25, 30 and 31 of novelty and an inventive step.